A Phase I Trial of AAV-Mediated Liver-Directed Gene Therapy for Hemophilia B

Stanford University Medical Center
The Children's Hospital of Philadelphia
Avigen, Inc.

Today's Focus

Issues Pertaining to IGLT of AAV Vector

SUMMARY OF AAV MEDIATED INTEGRATION IN LIVER

- ~5% of hepatocytes are stably modified with rAAV
- The proportion of integrated genomes is small, generally < 10% of double stranded vector DNA
- Gene expression from integrated and episomal AAV genomes parallels the proportion of vector DNA in each state
- There is no detectable increase in the proportion of integrated genomes over time
- The proportion of transduced cells with integrated genomes is small
- Most integrants are one or two copy genomes

Clinical Trial Objectives

- Test the hypothesis that AAVmediated liver-directed gene transfer is safe.
- Characterize the human immune response to the transgene product and to vector.
- Determine whether germline transmission of vector occurs following hepatic administration.
- Determine dose capable of producing clinically relevant Factor IX levels in the blood.

Clinical Trial Design

Phase I open-label, dose escalation safety trial of AAV-hFIX administration by infusion into the hepatic artery

Vector Administration

 Vector is infused into liver via a balloon occlusion catheter placed in the hepatic artery.

 Factor IX protein will be administered before and following procedure.

Subjects are observed at least 24 hours

Dose Escalation Plan

Group	Subjects	Dose/kg ¹ (v.g.) ²	Total Dose ³ (v.g.)	Observed F.IX levels in mice	Observed F.IX levels in dogs
1	2	2.0 x 10 ¹¹	1.4 x 10 ¹³	~1%	-
2	4	1.0 x 10 ¹²	7.0 x 10 ¹³	5%	4-12%4
3	4	5.0 x 10 ¹²	3.5 x 10 ¹⁴	20-30%	-

¹ dosing will be performed according to the patient weight obtained at the time of injection

² vector genomes

³ assuming a 70 kg adult

⁴ levels observed with a similar, but not identical construct.

New Exclusion Criteria*

- Patients unwilling to provide required semen
- of at least 20×10^6 sperm/mL, and motility (>50%). adequate semen volume (>1.5mL), sperm count Patients unable to provide semen samples of samples.

*Based on low-volume ejaculates in subject #2







Subject #1

- 63 Y/O male with severe factor IX deficiency
- S/P bilateral knee replacement 5 years PTA
- HIV negative
- HCV positive; HCV viral load negative
- Social Hx: father of 3; grandson with hemophilia

Subject #1

- Procedure (8/13/01)
- Received 2.0 x 10 ¹¹ vector genomes/kg
- No complications
- D/C home to referring HTC after 5 days

Subject # 1: Hematology Data

All CBCs within normal limits

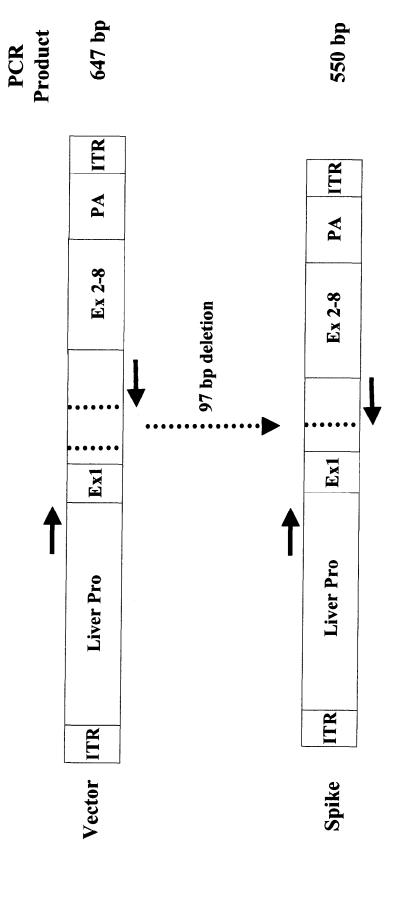
Subject # 1: LFT / PT

 All liver function tests and prothrombin time within normal limits

Subject #1: Coagulation Data

No therapeutic levels or inhibitors observed

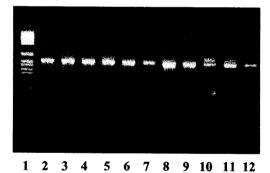
Design of Clinical PCR Assay



Example of PCR Analysis

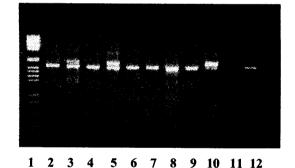
100 copies

BL D7



10 copies

BL D7



unspiked

BEL BEL D7 D7



1 2 3 4 5 6 7 8 9 10 11 12

Subject # 1: Detection of Vector Sequences by PCR

Subject # 1: Semen Analysis for Vector DNA

Samples performed in triplicate

* Week 3 sample fractionation: Motile sperm negative, seminal fluid positive, seminal fluid pellet positive, total semen positive.

Sensitivity of Assay: less than 1 copy per 30,000 haploid genomes.



Subject # 2

- 48 Y/O male with severe factor IX deficiency
- S/P bilateral knee replacement 1999, bilateral elbow replacement 2001
- HIV positive, HCV positive
- Hx Non-Hodgkin's lymphoma, 1986. Relapse in 1996
- Meds: Viracept, Epivir, Hivid.

Subject # 2

- Procedure (1/29/02)
- Received 2.0 x 10 ¹¹ vector genomes/kg
- No complications
- D/C to referring HTC after 7 days

Subject # 2: LFT / PT

All liver function tests and prothrombin time within normal limits

Subject # 2: Renal Function

Renal function within normal limits

Subject # 2: Hematology Data

All CBCs within normal limits

Subject # 2: Coagulation Data

- Factor IX levels?
- No inhibitors observed

^{*}Infusion with exogenous F.IX >14 days prior to FIX assay.

Subject # 2: Detection of Vector Sequence by PCR Analysis

Subject # 2: Semen Analysis for Vector DNA

*.07 mg of DNA extracted from month one sample

Conclusions

- Subjects #1 and #2 tolerated the procedure well
- Vector DNA present transiently in total semen from Subject #1, not present in motile sperm fraction at week 3

- Limited data in Subject #2
- Enrollment of subjects at mid-dose proceeds only if subject #2 shows absence of signal in motile sperm fraction